

SEQUENCE LISTING

<110> Terrett, Jonathan A

<120> NOVEL CANCER ASSOCIATED PROTEIN

<130> 2543-1-033

<150> PCT/GB02/02782

<151> 2002-06-14

<150> GB 0114643.0

<151> 2001-06-15

<150> GB 0205264.5

<151> 2002-03-06

<160> 10

<170> PatentIn version 3.1

<210> 1

<211> 310

<212> PRT

<213> Homo sapiens

<400> 1

Met Gly Asn Gln Val Glu Lys Leu Thr His Leu Ser Tyr Lys Glu Val
1 5 10 15

Pro Thr Ala Asp Pro Thr Gly Val Asp Arg Asp Asp Gly Pro Arg Ile
20 25 30

Gly Val Ser Tyr Ile Phe Ser Asn Asp Asp Glu Asp Val Glu Pro Gln
35 40 45

Pro Pro Pro Gln Gly Pro Asp Gly Gly Gly Leu Pro Asp Gly Gly Asp
50 55 60

Gly Pro Pro Pro Pro Gln Pro Gln Pro Tyr Asp Pro Arg Leu His Glu
65 70 75 80

Val Glu Cys Ser Val Phe Tyr Arg Asp Glu Cys Ile Tyr Gln Lys Ser
85 90 95

Phe Ala Pro Gly Ser Ala Ala Leu Ser Thr Tyr Thr Pro Glu Asn Leu
100 105 110

Leu Asn Lys Cys Lys Pro Gly Asp Leu Val Glu Phe Val Ser Gln Ala
115 120 125

Gln Tyr Pro His Trp Ala Val Tyr Val Gly Asn Phe Gln Val Val His
130 135 140

Leu His Arg Leu Glu Val Ile Asn Ser Phe Leu Thr Asp Ala Ser Gln
145 150 155 160

Gly Arg Arg Gly Arg Val Val Asn Asp Leu Tyr Arg Tyr Lys Pro Leu
165 170 175

Ser Ser Ser Ala Val Val Arg Asn Ala Leu Ala His Val Gly Ala Lys
180 185 190

Glu Arg Glu Leu Ser Trp Arg Asn Ser Glu Ser Phe Ala Ala Trp Cys
195 200 205

Arg Tyr Gly Lys Arg Glu Phe Lys Ile Gly Gly Glu Leu Arg Ile Gly
210 215 220

Lys Gln Pro Tyr Arg Leu Gln Ile Gln Leu Ser Ala Gln Arg Ser His
225 230 235 240

Thr Leu Glu Phe Gln Ser Leu Glu Asp Leu Ile Met Glu Lys Arg Arg
245 250 255

Asn Asp Gln Ile Gly Arg Ala Ala Val Leu Gln Glu Leu Ala Thr His
260 265 270

Leu His Pro Ala Glu Pro Glu Glu Gly Asp Ser Asn Val Ala Arg Thr
275 280 285

Thr Pro Pro Pro Gly Arg Pro Pro Ala Pro Ser Ser Glu Glu Glu Asp
290 295 300

Gly Glu Ala Val Ala His
305 310

<210> 2
<211> 1054
<212> DNA
<213> Homo sapiens

<400> 2
tgtgcaaagt accctggagt tggtttcgct ttctcccctt gcggcggtgt gaacgtgtgt 60
ccgcagcgtg atgggcaacc aggtggagaa attgaccac ctaagttaca aggaagttcc 120
cacggccgac ccgactggcg tggaccggga cgacgggccc cgcattgggg tctcctacat 180
tttctccaat gacgatgagg acgtggagcc gcagccgccg cctcaggggc cagatggcgg 240
cggcttgccc gacggtgggg acgggcccgc gccgcccag ccgcagccct acgatccgcg 300

```
<400> 5
ggctgctact gcaaacagtt cc 22
```

<210> 6
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer

<400> 6
ggtcaacgat ctgtaccgct ac 22

<210> 7
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer

<400> 7
gccgatcttg aactcgcgct tg 22

<210> 8
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer

<400> 8
ttcacctctc cgcgggtagc ct 22

<210> 9
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer

<400> 9
ggaagttacc cacatatacg gc 22

<210> 10
<211> 15
<212> PRT
<213> Homo sapiens

<400> 10

Ser Tyr Lys Glu Val Pro Thr Ala Asp Pro Thr Gly Val Asp Arg
1 5 10 15